



OIPE

ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/027,000

DATE: 05/13/2002

TIME: 09:51:58

Input Set : A:\GC696-SEQLIST.txt

Output Set: N:\CRF3\05102002\J027000.raw

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<110> APPLICANT: Dunn-Coleman, Nigel
 5
         Goedegebuur, Frits
 6
         Ward, Michael
 7
         Yao, Jian
 9
  <120> TITLE OF INVENTION: BGL4 Beta-Glucosidase and Nucleic Acids
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         Encoding the Same
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15 <141> CURRENT FILING DATE: 2001-12-18
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35 acggcgtaag agggaccaag ttetteaatg gegteeetge ggeetgette eettgeggea
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                                                                           900
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42 cgttcatgac ggcgtacaat ggcatcaatg gcgtgtcgtg cagcgagaac cctaaatatc
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46 teattgacea gagggetagg gaagttette agttegteaa gaagtgtget geeteeggag
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                                                                          1320
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49 gcaagaagaa gaagacgctg attgtcggcc ccaacgccaa gcaggccaca taccacggcg
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1620

RAW SEQUENCE LISTING DATE: 05/13/2002 PATENT APPLICATION: US/10/027,000 TIME: 09:51:58

Input Set : A:\GC696-SEQLIST.txt

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61 ttaacgccga	gteegtegee	azagagagag	ageaegaeea	catgaagete	cccaacatac	2100
61 Claacyccya	cigggagacc	gagggegeeg	accycycyay	catguagete	atcatacaga	2160
62 tggaccagct	cattgeegae	grayeegeeg	nagagagaga	caccegtcetc	gecaegeaga	2220
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84 Met Ala As 85 1 86 Glu Lys Va 87	p Ile Asp V 5 l Asp Leu L 20	eu Ala Gly	Ile Leu Lys 10 Ile Asp Phe 25	Trp His Thi	15 Lys Ala	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly	p Ile Asp V 5 1 Asp Leu L 20 s His Gly V	eu Ala Gly al Pro Ser	Ile Leu Lys 10 Ile Asp Phe 25	Trp His Thi	15 Lys Ala	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35	p Ile Asp V 5 1 Asp Leu L 20 s His Gly V	eu Ala Gly al Pro Ser 40	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe	Trp His Thr 30 Thr Asp Gly 45	15 Lys Ala Pro Asn	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar	p Ile Asp V 5 1 Asp Leu L 20 s His Gly V	eu Ala Gly (al Pro Ser 40 (ays Phe Phe	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala	15 Lys Ala Pro Asn	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr I	eu Ala Gly al Pro Ser 40 ys Phe Phe 55	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala	15 Lys Ala Pro Asn Cys Phe	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L	eu Ala Gly Cal Pro Ser 40 Ays Phe Phe 55 Aeu Gly Ser	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala	15 Lys Ala Pro Asn Cys Phe Leu Glu	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L	eu Ala Gly al Pro Ser 40 ys Phe Phe 55 eu Gly Ser	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Let	15 Lys Ala Pro Asn Cys Phe Leu Glu 80	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl	p Ile Asp V 5 1 Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L y Lys Met M	eu Ala Gly al Pro Ser 40 ys Phe Phe 55 eu Gly Ser	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Let	15 C Lys Ala V Pro Asn Cys Phe Leu Glu 80 C Ala His	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L 7 y Lys Met M 85	eu Ala Gly al Pro Ser 40 ys Phe Phe 55 eu Gly Ser 0 let Gly Lys	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Sen	15 C Lys Ala V Pro Asn Cys Phe Leu Glu 80 C Ala His 95	
84 Met Ala As 85 1 86 Glu Lys Va 87	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L y Lys Met M 85 u Gly Pro I	eu Ala Gly al Pro Ser 40 ys Phe Phe 55 eu Gly Ser 0 let Gly Lys	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser	15 C Lys Ala Y Pro Asn Cys Phe Leu Glu 80 C Ala His 95 L Gly Gly	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L 7 y Lys Met M 85 u Gly Pro T	Tal Pro Ser 40 Tys Phe Phe 55 Theu Gly Ser 0 The Gly Lys The Ile Asn	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu	15 C Lys Ala V Pro Asn Cys Phe Leu Glu 80 C Ala His 95 I Gly Gly	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le 97 98 Arg Gly Ph	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L 7 y Lys Met M 85 u Gly Pro T 100 e Glu Ser I	deu Ala Gly Tal Pro Ser 40 Tys Phe Phe 55 Teu Gly Ser 0 Tet Gly Lys Thr Ile Asn The Gly Glu	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu 110 Leu Ala Gly	15 C Lys Ala V Pro Asn Cys Phe Leu Glu 80 C Ala His 95 I Gly Gly	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le 97 98 Arg Gly Ph 99 11	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L 7 y Lys Met M 85 u Gly Pro T 100 e Glu Ser I	deu Ala Gly Tal Pro Ser 40 Tys Phe Phe 55 Tys Gly Ser 0 Tet Gly Lys Thr Ile Asn Tle Gly Glu 120	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg 105 Asp Pro Phe	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu 110 Leu Ala Gly 125	15 C Lys Ala Y Pro Asn Cys Phe Leu Glu 80 C Ala His 95 G Gly Gly Y Leu Gly	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le 97 98 Arg Gly Ph 99 11 100 Ala Ala A	p Ile Asp V 5 l Asp Leu L 20 s His Gly V g Gly Thr L y Thr Ser L 7 y Lys Met M 85 u Gly Pro T 100 e Glu Ser I	eu Ala Gly Cal Pro Ser 40 Eys Phe Phe 55 Eeu Gly Ser 0 Eet Gly Lys Chr Ile Asn Cle Gly Glu 120 Arg Gly Ile	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg 105 Asp Pro Phe	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu 110 Leu Ala Gly 125 r Gly Val Gl	15 C Lys Ala Y Pro Asn Cys Phe Leu Glu 80 C Ala His 95 G Gly Gly Y Leu Gly	
84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le 97 98 Arg Gly Ph 99 11 100 Ala Ala A 101 130	p Ile Asp V 5 l Asp Leu I 20 s His Gly V g Gly Thr I y Thr Ser I 7 y Lys Met M 85 u Gly Pro T 100 e Glu Ser I 5 la Leu Ile	eu Ala Gly Cal Pro Ser 40 Eys Phe Phe 55 Eeu Gly Ser 0 Eet Gly Lys Chr Ile Asn 120 Arg Gly Ile 135	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg 105 Asp Pro Phe	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu 110 Leu Ala Gly 125 r Gly Val Gl	15 Lys Ala Pro Asn Cys Phe Leu Glu 80 Ala His 95 Gly Gly Y Leu Gly In Ala Thr	
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84 Met Ala As 85 1 86 Glu Lys Va 87 88 Leu Pro Ly 89 35 90 Gly Val Ar 91 50 92 Pro Cys Gl 93 65 94 Glu Ala Gl 95 96 Val Ile Le 97 98 Arg Gly Ph 99 11 100 Ala Ala A 101 130	p Ile Asp V 5 l Asp Leu I 20 s His Gly V g Gly Thr I y Thr Ser I 7 y Lys Met M 85 u Gly Pro I 100 e Glu Ser I 5 la Leu Ile is Phe Leu	eu Ala Gly al Pro Ser 40 ys Phe Phe 55 eu Gly Ser 0 et Gly Lys chr Ile Asn 120 Arg Gly Ile 135 Cys Asn Asp	Ile Leu Lys 10 Ile Asp Phe 25 Leu Arg Phe Asn Gly Val Thr Phe Asn 75 Glu Ala Ile 90 Met Gln Arg 105 Asp Pro Phe e Gln Ser Th	Trp His Thr 30 Thr Asp Gly 45 Pro Ala Ala 60 Gln Thr Leu Ala Lys Ser Ser Pro Leu 110 Leu Ala Gly 125 r Gly Val Gl 140 p Arg Arg Me	15 Lys Ala Pro Asn Cys Phe Leu Glu 80 Ala His 95 Gly Gly V Leu Gly Leu Gly In Ala Thr	

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Input Set : A:\GC696-SEQLIST.txt

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114	Asp	Leu	Glu	Met	Pro	Gly	Pro	${\tt Pro}$	Arg	Phe	Arg	Gly	Glu	Thr	Leu	Lys
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116	Phe	Asn	Val	Ser	Asn	Gly	Lys	${\tt Pro}$	Phe	Ile	His	Val	Ile	Asp	Gln	Arg
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124	Glu	Asn	Asn	Val	Leu	Pro	Leu	Ser	Lys	Lys	Lys	Lys	Thr	Leu	TTE	var
125					325	_			_	330	-1	a 1	a 1	a	335	110
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	Leu	Arg			Tyr	Ala	Val		Pro	Pne	Asp	GIY	365	Ser	гуз	GIII
129			355	_	_	~	m	360	77. 7	01	7 J -	m		Thr	Wa l	Dro
	Leu		Thr	Pro	Pro	ser		Tnr	vaı	СТУ	Ата	380	1111	Thr	vaı	FIO
131	_	370		01	01	01 -	375	T 011	mh∽	Dro	A an		λla	Dro	Glv	Met
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		ASP	GIU	420		FIIC	1111	цуз	425	115 _P	rice	1110		430	E	-1-
137	m	III a	Dro			λla	Δen	Thr		Tvr	Δla	Asp	Met		Glv	Thr
		HIS	435		АТа	АТа	изр	440	115	- 1 -	mil	тор	445	0	-1	
139	Пттх	Thr	A J S	λen	Glu	Δen	Cvs	Thr	Ψvr	Glu	Leu	Glv	Leu	Ϋal	Val	Cys
141		450		кар	Giu	пор	455	1 111	-1-	0		460				-
141	G1v	Thr	Δla	Lvs	Δla	Ψvr		Asp	Asp	Gln	Leu		Val	Asp	Asn	Ala
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145		цуз	OIL	,,,,	485					490	. 4				495	
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Input Set : A:\GC696-SEQLIST.txt

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	Gln		Trp	Tvr	Glv	Glv		Glu	Thr	Glv	Asn		Ile	Ala	Asp	Val	
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171		690					695					700					
		Thr	Thr	Phe	Ala		Ser	Asn	Leu	Ser		Ser	His	Lys	Asp		
	705	T 011	Com	37 - 1	Cor	710	Cor	นาไ	T 17.0	λαn	715	C1 17	cor	Wa 1	Pro	720	
175	гуѕ	ьeu	ser	vaı	725	Leu	ser	Val	цуѕ	730	1111	СТУ	SET	vai	735	Gry	
	Ala	Gln	Val	Ala		Leu	Tyr	Val	Lys		Leu	Gln	Ala	Ala	Lys	Ile	
177				740			-		745					750	_		
		Arg		Val	Lys	Glu	Leu	_	Gly	Phe	Ala	Lys		Glu	Leu	Gln	
179		a 1	755	m1	.		*** 1	760	T1.	01	a 1	01 =	765	T	m	wa 1	
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			Tvr	Phe	Asp	Glu		Ara	Asp	Gln	Trp		Val	Glu	Lys	Glv	
	785		-1-		F	790		5			795	-1-			- 4 -	800	
184	Asp	Tyr	Glu	Val	Ile	Val	Ser	Asp	Ser	Ser	Ala	Ala	Lys	Asp	Gly	Val	
185					805	_	_		_	810	_		_		815		
		Leu	Arg		Lys	Phe	Thr	Val		Glu	Thr	Tyr	Trp		Ser	Gly	
187	Val			820					825					830			
		0> SI	EQ II	о мо	: 3												
			ENGTI													•	
			YPE:														
			RGAN:			chode	erma	ree	sei								
			EQUE			~~ ~				~ ~ ~ ~	a+ a a	0001	- ~ ~ ~	202	~ ~ ~ <i>~</i>	rt.aaat	60
																gtcgat ccctct	120
																gteect	180
																ctcgaa	240
201	gag	gcag	gta a	agate	gatg	gg ca	aaaga	aggc	c ato	cgcta	aaga	gtg	cgcat	tgt (gatco	ctcggc	300
	_					-			-		-	_				ggtgag	360
																actgga	420 480
																atggtg attgct	540
203	cage	ayca	ccy (Louc	guy	-9 9 <u>'</u>	9000	ccg	- yu		cucy	Caci		, ·	coage	900	240

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/027,000

DATE: 05/13/2002 TIME: 09:51:59

Input Set : A:\GC696-SEQLIST.txt
Output Set: N:\CRF3\05102002\J027000.raw

206	gtgcgagact	cccagccggg	tgcgttcatg	acggcgtaca	atggcatcaa	tggcgtgtcg	600
				atgcttcgaa			660
208	ctaatcatga	gcgactggta	cggcacatac	agtaccacag	aagccgttgt	ggcaggcctc	720
209	gacctcgaga	tgcccggacc	tccacgcttc	cgaggagaaa	cactcaagtt	caacgtctcc	780
210	aacggaaagc	cctttatcca	cgtcattgac	cagagggcta	gggaagttct	tcagttcgtc	840
211	aagaagtgtg	ctgcctccgg	agtgacggag	aacggccccg	agacgactgt	caacaacacc	900
				ggcaacgagg			960
				aagaagacgc			1020
				gccgcactca			1080
215	ccctttgacg	gcctcagcaa	gcagctcgag	acgccgccat	cgtacaccgt	cggcgcctac	1140
				tgcctcacgc			1200
				acccctaacc			1260
				gactactacc			1320
				gccgacgagg			1380
				gtagacgacc			1440
				ggctccgcca			1500
				ttcaagatcg			1560
				ggccacggct			1620
				aagtccgtcg			1680
				gactgggaga			1740
				ctcattgccg			1800
				cccgaggaga			1860
				aacgagacgg			1920
				ctgtccctca			1980
				gaggccgggc			2040
				gccgacaagg			2100
				tccaatctct			2160
				accggctccg			2220
				aagattaacc			2280
				gagacgaagg			2340
				gagcgggatc			2400
				gcagcgaagg		gctcaggggt	2460
238	aagtttacgg	tgggagagac	gtattggtgg	tctggcgtgt	aa		2502

VERIFICATION SUMMARY

DATE: 05/13/2002

PATENT APPLICATION: US/10/027,000

TIME: 09:52:00

Input Set : A:\GC696-SEQLIST.txt